# Effects of Perinatal PBDE Exposure on Hepatic Phase I, Phase II, Phase III, and Deiodinase 1 Gene Expression Involved in Thyroid Hormone Metabolism in Male Rat Pups

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Previous studies demonstrated that perinatal exposure to polybrominated diphenyl ethers (PBDEs), a major class of brominated flame retardants, may affect thyroid hormone (TH) concentrations by inducing hepatic uridinediphosphateglucoronosyltransferases (UGTs). This study further examines effects of the commercial penta mixture, DE-71, on genes related to TH metabolism at different developmental time points in male rats. DE-71 is predominately composed of PBDE congeners 47, 99, 100, 153, 154 with low levels of brominated dioxin and dibenzofuran contaminants. Pregnant Long-Evans rats were orally administered 1.7 (low), 10.2 (mid), or 30.6 (high) mg/kg/ day of DE-71 in corn oil from gestational day (GD) 6 to postnatal day (PND) 21. Serum and liver were collected from male pups at PND 4, 21, and 60. Total serum thyroxine (T<sub>4</sub>) decreased to 57% (mid) and 51% (high) on PND 4, and 46% (mid) dose and 25% (high) on PND 21. Cyp1a1, Cyp2b1/2, and Cyp3a1 enzyme and mRNA expression, regulated by aryl hydrocarbon receptor, constitutive androstane receptor, and pregnane xenobiotic receptor, respectively, increased in a dose-dependent manner. UGT-T<sub>4</sub> enzymatic activity significantly increased, whereas age and dosedependent effects were observed for Ugt1a6, 1a7, and 2b mRNA. Sult1b1 mRNA expression increased, whereas that of transthyretin (Ttr) decreased as did both the deiodinase I (D1) enzyme activity and mRNA expression. Hepatic efflux transporters Mdr1 (multidrug resistance), Mrp2 (multidrug resistance-associated protein), and Mrp3 and influx transporter Oatp1a4 mRNA expression increased. In this study the most sensitive responses to PBDEs following DE-71 exposure were CYP2B and D1 activities and Cyb2b1/2, d1, Mdr1, Mrp2, and Mrp3 gene expression. All responses were reversible by PND 60. In conclusion, deiodination, active transport, and sulfation, in addition to glucuronidation, may be involved in disruption of TH homeostasis due to perinatal exposure to DE-71 in male rat offspring.

Key Words: PBDE; flame retardant; thyroid; hepatic metabolism; rat.

Brominated flame retardants (BFRs) are added to consumer products to delay the ignition and burning of materials, thereby saving lives and reducing property damage. Polybrominated diphenyl ethers (PBDEs), a major type of BFRs, consist of a family of 209 possible congeners; however, what is actually used commercially is a flame-retardant mixture. Three PBDE commercial flame-retardant products have been produced which are mixtures of congeners, and referred to based on the average level of bromination in the mixture: decabrominated diphenyl ether (e.g., DE-83R); octabrominated diphenyl ether, for example, (DE-79); and pentabrominated diphenyl ether (e.g., DE-71). DE-71 is used in polyurethane foams, upholstery, car and airline seats, office and household furniture, carpet pads, mattresses, and pillows (Birnbaum et al., 2004). DE-71 consists primarily of the congeners BDE 47, 99, 100, 153, 154 (LaA Guardia et al., 2006) and has been found to contain varying low levels of polybrominated dibenzop-dioxins and dibenzofurans (Hanari et al., 2006). This commercial penta mixture was used as an additive flame retardant which allows it to escape into the environment (Kim et al., 2006) including household dust (Stapleton et al., 2005). The European Union has banned production and use of all PBDE mixtures. Although, the sole U.S. producer voluntarily stopped production of DE-71 (penta) and DE-79 (octa) at the end of 2004 and 11 states have banned their production, use and import, the presence of PBDEs in the environment and biota continues. The persistence of PBDEs is attributed to its lipophilic properties and bioaccumulative nature.

PBDEs have been detected in human serum, breast milk, and adipose tissue at varying levels (Birnbaum *et al.*, 2004; McDonald, 2002). PBDEs, as polychlorinated biphenyls (PCBs) and dioxins, are structurally similar to thyroid hormones (THs) and therefore, may act as endocrine disruptors via alterations in TH homeostasis (Darnerud *et al.*, 2001; Hooper and McDonald, 2000; Kuriyama *et al.*, 2007; Tseng *et al.*, 2008). As reviewed by Costa and Giordano (2007), PBDEs have also been identified as developmental neurotoxicants. A single low exposure of BDE 47, 99, and 209 to

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mice during neonatal development permanently alters behavior (Eriksson *et al.*, 2001; Rice *et al.*, 2007; Viberg *et al.*, 2003).

TH has a crucial role in fetal brain development in both animals and humans. During fetal and early neonatal periods, disorders of TH homeostasis may result in motor and cognitive disorders (Sher et al., 1998). The thyroid axis contains numerous sites in which xenobiotics can alter the hormonal balance. Decreases in TH concentrations are often observed in rodents with increases in hepatic thyroxine (T<sub>4</sub>) glucuronidation leading to increased biliary elimination of the conjugated hormone (Barter and Klaassen, 1992; Vansell and Klaassen, 2002). Maternal administration of DE-71 during gestation and lactation results in a reduction and functional disruption of pup serum T<sub>4</sub> levels with increase in uridinediphosphateglucoronosyltransferases (UGTs) activity in rats (Ellis-Hutchings et al., 2006; Zhou et al., 2001). Adult rats exposed to PBDEs also have reduced concentration of circulating T<sub>4</sub> which is associated with an induction in hepatic UGTs (Hallgren and Darnerud, 2002). A 14-day exposure to BDE 47 at 18 mg/kg/day reduced total serum T<sub>4</sub> concentrations in rats (Darnerud and Sinjari, 1996). Additionally, a 4-day exposure in female mice resulted in 50% decrease in circulating total T<sub>4</sub> concentrations at 100 mg/kg/day of BDE 47 (Richardson et al., 2008). Collectively, these studies demonstrate that DE-71 and individual PBDE congeners alter TH homeostasis in developing and adult rodents.

Although it has been suggested that decreases in circulating THs may be due to induction of UGT, it is unclear if this alone is responsible for the decrease in circulating TH. UGT1Adeficient Gunn rats exposed to phenobarbital (PB) or PCBs had decreases in serum total T4 demonstrating a lack of total dependence on glucuronidation (Collins and Capen, 1980; Kato et al., 2004). Sulfotransferases (SULTs) also play a key role in TH homeostasis, as they are the major pathway for T<sub>4</sub> conjugation in humans (Visser, 1994). THs regulate SULT gene expression in an isoform-specific manner (Dunn and Klaassen, 2000). Studies using adult rat hepatocytes demonstrated SULT1B1 and 1C1 conjugate iodothyronines (Kester et al., 2003). Furthermore, sulfation increases the degradation of T<sub>4</sub> by type I iodothyronine deiodinase (D1) (Kaptein et al., 1997). D1 is a microsomal selenoenzyme which catalyzes deiodination of the prohormone T<sub>4</sub> to the active 3,5,3'triiodothyronine (T<sub>3</sub>) as well as to the inactive metabolite reverse-T<sub>3</sub> (rT<sub>3</sub>), and subsequently to the inactive metabolite 3,3'-diiodothyronine (T<sub>2</sub>) (Visser, 1994). These studies suggest sulfation and deiodination, along with glucuronidation, may play a role in altering the homeostasis of TH (Hood and Klaassen, 2000). Further, in vivo studies show that PCB metabolites (Kato et al., 2004) and hydroxylated PBDEs (Meerts et al, 2000) can bind to transthyretin (Ttr), a major TH transport protein in plasma, which also may cause a decrease in serum total T4.

Phase I, II, and III xenobiotic metabolizing enzymes (XMEs) play fundamental roles in metabolism and elimination of xenobiotics. XMEs can be present either at the basal level and/

or are induced or inhibited after exposure to xenobiotics via activation of a variety of nuclear receptors including aryl hydrocarbon receptor (AhR), constitutive androstane receptor (CAR), and pregnane xenobiotic receptor (PXR) (Tompkins and Wallace, 2007; Xu et al., 2005). Recent studies show that BDE 47 predominantly activates hepatic CAR and to a lesser extent PXR regulated genes following a 4-day exposure in adult mice (Pacyniak et al., 2007; Richardson et al., 2008). The role of UGTs in the DE-71 and individual PBDE-mediated decrease in TH has been examined in the adult and developing rodent, however, the role of other phase II metabolizing enzymes (i.e., SULTs) and phase III transporters in the reduction of TH has yet to be examined. A correlation between induction of hepatic UGT mRNAs, multidrug resistanceassociated protein (Mrp2) mRNA, and organic anion transporting protein (Oatp1a4) mRNA levels, with decreases in serum TH concentrations has been observed (Hagenbuch, 2007; Mitchell et al., 2005; Ribeiro et al., 1996). P-glycoprotein (Pgp), a product of the multidrug resistance (Mdr) gene, is expressed in various tissues and plays an important role in drug absorption and disposition, whereas alterations in the expression levels of Pgp are involved in the variability of pharmacokinetics of many drugs and xenobiotics. There is evidence that multidrug-resistance transporters are important in unconjugated TH efflux (Mitchell et al., 2005). In a recent study, BDE-47 altered Mdr1a and Mrp3 mRNA expression levels in adult mice (Richardson et al., 2008). Studies using Xenopus laevis oocytes identified transporters involved in TH uptake; Na(+) taurocholate cotransporting polypeptide (NTCP), and the organic anion transporting polypeptide (OATP) (Friesema et al., 2005). Collectively, these studies indicate that active transport, in addition to sulfation, deiodination, and altered serum binding, could also play a role in the disposition of TH.

This study examines how perinatal exposures to DE-71 alter the expression of hepatic genes involved in TH homeostasis in Long-Evans male rat pups in order to further the risk assessment of this commercial penta-BDE mixture. In addition to cytochrome P450s (CYPs) and UGTs mRNA expression and enzyme activity, SULTs, D1, and transporters were also examined to determine other possible mechanisms involved in disruption of TH homeostasis following early life exposure to DE-71. The basal expression of major XMEs and transporters in the developing rat pups is also described (Supplementary Data).

# METHODS AND MATERIALS

**Animals.** Time-pregnant Long-Evans rats, approximately 80–90 days of age, were obtained from Charles River Laboratories, Inc. (Raleigh, NC) on GD 2, and allowed 4 days acclimation in an American Association for Accreditation of Laboratory Animal Care–approved animal facility prior to being treated. Dams were housed individually in plastic cages ( $45 \times 24 \times 20$  cm) with sterilized pine shavings as bedding, which was changed twice a week except on the day of parturition (i.e., GD 21). Rats were provided with Purina 5001 Rodent Chow (Ralston Purina Co., St Louis, MO) and tap water

ad libitum. On PND 21, offspring were counted, sexed, and group-weighed by sex. Average pup weight by sex was calculated by dividing the group weight by the number of pups. Pups were weaned on PND 21, and housed by gender in groups of two per cage.

Chemicals and treatment. DE-71 (penta-BDE, lot 7550OK20A) was generously supplied by the Great Lakes Chemical Corporation (West Lafayette, IN). DE-71 is a mixture that consists primarily of tetra and penta congeners. The stock DE-71 solution (300 mg/ml) was prepared by mixing the compound with corn oil and sonicating for 30 min at 40°C. The desired dosing solutions were obtained by serial dilution with corn oil. The dams were orally dosed (2 ml/kg), via gavage, with DE-71 (0, 1.7, 10.2, or 30.6 mg/kg/day) from GD 6 through PND 21, except for PND 0 (day of birth) when dams were left undisturbed. On PND 4, litters were culled to eight pups per litter, four males and four females. Pups were euthanized on PND 4, 21, or 60 with CO2 asphyxiation followed by exsanguinations via cardiac puncture where blood and liver were collected. Liver samples were removed immediately and frozen in liquid nitrogen. Serum was obtained after clotting whole blood on ice, followed by centrifugation at 12000 g at 4°C for 20 min. All serum and liver samples for each age point were obtained from a minimum of eight litters, and were stored at -80°C until analysis for TH (T<sub>4</sub>, T<sub>3</sub>) concentrations, hepatic enzyme activity, and mRNA expression analysis. The liver was the experimental unit.

*TH assay.* Serum concentrations of total T<sub>4</sub> and T<sub>3</sub> were measured in duplicate as previously described (Goldey *et al.*, 1995) by using standard radioimmunoassay kits (Diagnostic Products Corp., Los Angeles, CA).

Hepatic enzyme activity assay. Liver microsomal fractions were prepared as described previously (DeVito et al., 1993). Microsomal protein concentrations were determined using the Bio-Rad protein assay kit (Bio-Rad, Richmond, CA) with bovine serum albumin (BSA) (Sigma, St Louis, MO) as the standard. Hepatic microsomal ethoxyresorufin-O-dealkylase (EROD) activity (a marker for CYP1A1 activity), pentoxyresorufin-O-dealkylase (PROD) activity (a marker of CYP2B activity), and benzyloxyresorufin-O-dealkylase (BROD) activity (a marker of CYP3A activity) were assayed according to DeVito et al. (1993). All substrate concentrations were 1.5nM. EROD, PROD, and BROD values were calculated as pmol resorufin per mg protein per min.

*UGT-T<sub>4</sub> activity assay.* UGT-T<sub>4</sub> activity was determined by using a modified assay (Zhou *et al.*, 2002) based on a previously published method (Visser *et al.*, 1993). A 100-μl aliquot of microsomes (2 mg protein per 100mM Tris/HCl buffer pH 7.4) was incubated at 37°C with purified <sup>125</sup>I-T<sub>4</sub> (NEN Life Science Products, Inc., Boston, MA), 6-n-propyl-2-thiouracil (PTU), and uridine diphosphoglucuronic acid (UDPGA) (or no UDPGA for blank) (Sigma) over a 30-min period. The reaction was stopped with methanol (Sigma) followed by centrifugation and mixing the supernatant with 0.1M HCl (Sigma). The formed glucuronyl T<sub>4</sub> was separated by chromatography on 2 ml of lipophilic sephadex (Sigma) LH-20 columns (Superlco, Bellefonte, PA) and radioactivity determined by gamma scintillation spectrometry.

Outer-ring deiodinase activity. D1 was determined by quantifying the amount of  $I^{125}$  released from  $I^{125}\text{-r}T3$  (NEN) (Leonard and Rosenberg, 1980). All reactions contained 75  $\mu l$  of 0.1M phosphate (pH 7.0) and 2mM ethylenediaminetetraacetic acid (EDTA) buffer (Sigma), 25  $\mu l$  of 2 mg/ml microsomal protein, and 1mM PTU (Sigma) (in blanks to determine spontaneous deiodination). The reaction was stopped by adding 50  $\mu l$  of 4% BSA (Sigma) in 1mM PTU solution. Ten percent trichloroacetic acid (TCA; Sigma) was added, centrifuged, and the supernatant was loaded on 2 ml of lipophilic sephadex (Sigma) in a LH-20 column (Superlco) (Rooda et~al., 1987). Free iodine was eluted using 0.1M HCl (Sigma) and radioactivity determined as above.

SULT activity. Iodothyronine SULT activities were assayed by incubation of  $1\mu M$  of  $T_4$ , and  $10^5$  cpm of  $^{125}$ I-labeled  $T_4$  (NEN) for 30 min at 37°C with 20–40  $\mu g$  protein/ml of liver cytosol in the presence or absence (blank) of

 $50\mu M$  3'-phosphoadenosine 5'-phosphosulfate (PAPS; Sigma) in 0.2 ml of 0.1M phosphate (pH 7.2) and 2mM EDTA (Sigma) (Kaptein  $\it{et~al.}$ , 1997). The mixtures were applied to 1 ml of lipophilic sephadex (Sigma) LH-20 minicolumns (Superlco), and equilibrated in 0.1M HCl (Sigma). Iodine, sulfated iodothyronines, and nonsulfated iodothyronines were successively eluted with  $2\times 1$  ml of 0.1M HCl,  $6\times 1$  ml of ethanol/water (20/80, vol/vol), and  $3\times 1$  ml of ethanol/0.1M NaOH (50:50, vol/vol) (Sigma), respectively. Fractions were collected and measured for radioactivity.

RNA isolation and real-time RT-PCR analysis. Total RNA was isolated using the RNeasy Midi Kit with DNase I digestion performed during column purification (Qiagen, Hilden, Germany). To assess the integrity of the RNA samples, the 2100 Bioanalyzer was used (Agilent Technologies, Palo Alto, CA) with RNA Integrity Numbers (RINs) greater than 8.1. Samples with RINs less than 8.1 were not analyzed by RT-PCR (n = 5 per group). Real-time RT-PCR was performed using the ABI Prism 7700 Sequence Detection System (ABI, Foster City, CA). One-hundred nanograms of total RNA was used for each reaction. cDNA was synthesized using TaqMan Reverse Transcriptase Kits (ABI, Foster City, CA). PCR was then performed on all cDNAs using TaqMan Universal PCR Master Mix and TaqMan Gene Expression Assays (ABI) unless otherwise indicated. The probe/primer sets are Cyp1a1 (Rn00487218\_m1), Cyp3a1 (Rn01412959\_g1), Ugt1a1 (Rn007549447\_m1), Ugt1a6 (Rn00756113\_m1), Ugt1a7 (Rn01643133\_g1), Ugt2b (Rn0234965\_m1), Sult1a1 (Rn01510633\_ m1), Sult1b1 (Rn01524591\_g1), Sult1c1 (Rn01528019\_m1) Mdr1(Rn01529259\_ m1), Mrp2 (Rn00680936\_m1), Mrp3 (Rn00589786\_m1), Oatp1a4 (Rn00584891\_m1), Oat2 (Rn00585513\_m1), Ntcp (Rn00690543\_m1), Dio1 (Rn00562124\_m1), Ttr (Rn01416940\_m1), Cyp2b1 (Fw-5'GTGGGCCA-AGCTGAGGAT3', Rev-5'GAGAATCGCCGAAGGGC3', Probe-5'ATCG-CTGTGATTGAGCCAATCTTCAAGG3') and Cyp2b2 (Fw-TTCTGCG-CATGGAGAAAGTG3', Rev-5'GATCATGAGGTTCTCATGATGGAA3', Probe-5'CCTGCATGGATGAGAGAGGAGAGAGTCG3'). The thermal cycle condition for the RT reaction was as follows: 10 min at 25°C, 30 min at 48°C, and then 5 min at 95°C. The PCR reaction was performed as follows: 2 min at 50°C (uracil-DNA glycosylase activation), 10 min at 95°C (activation of Taq DNA polymerase), and 40 cycles of denaturation at 95°C for 15 s followed by annealing and extension at 60°C for 1 min. Data were analyzed using the Sequence Detection Systems software (ABI). All RT-PCR data were quantified by the  $\Delta\Delta C_t$  method (Applied Biosystems User Bulletin 2). These data were compared with a calibrator sample (a test sample where all other samples are relative expression levels of it) and an endogenous control (18S). In the  $\Delta\Delta C_t$ method, the control group is used as a reference point for all other dose groups; therefore, all samples are expressed as fold difference as compared with control.

**Data analysis.** The statistical intergroup comparisons of  $T_4$ ,  $T_3$ , RT-PCR, and enzyme activity data were determined by using a two-way ANOVA (examining the effect of dose, age, and their interaction on response) with a follow up contrast of dose to control in each age group. The follow up contrast were Bonferroni adjusted. The levels of probability of statistical significance are p < 0.05 and 0.01.

# RESULTS

Serum  $T_4$  and  $T_3$  Levels

A significant interaction between age and dose ( $F_{6,48}$  = 21.56; p < 0.001) was observed for total serum  $T_4$  (Table 1). A *post hoc* test indicated no significant effects at the low dose for any time points tested. However, at PND 4, exposure to DE-71 decreased total serum  $T_4$  to 57 and 51% of control levels with exposure to 10.2 and 30.6 mg/kg/day, respectively. There was also a decrease of total serum  $T_4$  at PND 21 with levels 46 and 25% of controls with perinatal exposure to 10.2 and 30.6 mg/kg/day, respectively. By PND 60, no differences

TABLE 1
Circulating Total T<sub>4</sub> and T<sub>3</sub> Levels (ng/ml) in Male Pups during
Perinatal Exposure to a Penta PBDE Mixture, DE-71

		PND			
DE-71	4	21	60		
Total T <sup>4</sup> levels					
0 mg/kg/day	$17.15 \pm 0.58$	$53.18 \pm 2.57$	$71.14 \pm 4.68$		
1.7 mg/kg/day	$18.73 \pm 1.74$	$47.68 \pm 3.11$	$67.96 \pm 7.73$		
10.2 mg/kg/day	$9.80 \pm 0.52*$	24.68 ± 1.26*	$72.05 \pm 11.21$		
30.6 mg/kg/day	$8.46 \pm 0.55$ *	$13.64 \pm 0.93*$	$68.13 \pm 5.98$		
Total T <sup>3</sup> levels					
0 mg/kg/day	$0.26 \pm 0.01$	$1.08 \pm 0.06$	$0.90 \pm 0.05$		
1.7 mg/kg/day	$0.28 \pm 0.02$	$1.14 \pm 0.05$	$0.76 \pm 0.05$		
10.2 mg/kg/day	$0.27 \pm 0.01$	$0.90 \pm 0.04$	$0.77 \pm 0.05$		
30.6 mg/kg/day	$0.29 \pm 0.01$	$0.88 \pm 0.04$	$0.83 \pm 0.05$		

*Note*.  $T_4$  and  $T_3$  serum levels were analyzed from male rats at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control: \*p < 0.05. Data are expressed as mean  $\pm$  SD, n = 7-9 rats.

between control and dose groups were seen. Interestingly, there was no significant effect of dose on serum total  $T_3$  levels at any age group. In this study, decreases in total  $T_4$  levels were similar in both males and females. Therefore, we selected males only for understanding the mechanism as female offspring from this cohort were used for reproductive and neurotoxicological endpoints (Kodavanti *et al.*, unpublished data).

Cytochrome P450s: Enzyme Activity and Gene Expression

Cyp1a1 mRNA expression and EROD activity are used as markers for Ah receptor activation (Table 2). There was a significant interaction of age and dose ( $F_{6.48} = 37.16$ ; p <0.0001) for Cyp1a1. A post hoc test indicates an effect at all doses tested at PND 4 and PND 21, and only for 30.6 mg/kg/ day at PND 60. Cyp1a1 mRNA expression was induced at the 1.7, 10.2, and 30.6 mg/kg/day treatment doses by 16.1-, 44.3-, 154.2-fold at PND 4, and 4.6-, 10.2-, and 20.1-fold at PND 21, respectively. There remained a residual 5.6-fold increase in Cyp1a1 mRNA expression with the 30.6 mg/kg/day treatment group at PND 60. In agreement with this, EROD indicated a significant interaction of age and dose ( $F_{6.48} = 152.3$ ; p <0.0001) at all doses tested for PND 4 and PND 21, but not at PND 60. Hepatic EROD increased at the 1.7, 10.2, and 30.6 mg/kg/day doses by 23.3-, 49.5-, and 215.3-fold at PND 4, and 4.1-, 17.1-, and 31.9-fold at PND 21, respectively.

Cyp2b1 and Cyp2b2 mRNA expression and PROD activity are used as markers for CAR receptor activation (Maglich *et al.*, 2002; Waxman, 1999; Yamada *et al.*, 2006). There was a significant interaction of age and dose for Cyp2b1 ( $F_{6,48} = 33.23$ ; p < 0.0001) and for Cyp2b2 ( $F_{6,48} = 41.28$ ; p < 0.0001). A *post hoc* test indicated an effect at all doses tested at PND 4 and 21 but not at PND 60. Hepatic Cyp2b1 mRNA

TABLE 2
Effect of DE-71 on Hepatic Cytochrome P450 Gene Expression and Protein Activity

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
Cyp1a1 <sup>a</sup>	0	$1.6 \pm 0.2$	$8.2 \pm 2.5$	$0.02 \pm 0.0$
• •	1.7	26.0 ± 9.9*	38.1 ± 11.6**	$0.02 \pm 0.0$
	10.2	71.3 ± 19.7*	84.1 ± 24.8*	$0.04 \pm 0.0$
	30.6	248.3 ± 29*	165.8 ± 55.6*	0.10 ± 0.1**
$EROD^b$	0	$4.2 \pm 1.1$	$108.9 \pm 12$	$87.2 \pm 9.0$
	1.7	98.2 ± 13.4**	443.2 ± 50.4**	$84.3 \pm 7.2$
	10.2	208.1 ± 62.3**	1872.0 ± 203.2**	$83.2 \pm 5.1$
	30.6	904.1 ± 113.2**	3472.2 ± 439.1**	$85.7 \pm 6.8$
Cyp2b1 <sup>a</sup>	0	$1.7 \pm 0.1$	$5.5 \pm 0.8$	$0.3 \pm 0.0$
71	1.7	6.6 ± 2.7**	37.7 ± 17.3**	$0.4 \pm 0.2$
	10.2	21.6 ± 7.4**	89.7 ± 33.2*	$0.5 \pm 0.1$
	30.6	$33.7 \pm 6.8*$	147.7 ± 20.1*	$0.5 \pm 0.1$
Cyp2b2 <sup>a</sup>	0	$1.7 \pm 0.3$	$6.5 \pm 1.5$	$0.7 \pm 0.1$
<b>71</b>	1.7	$3.4 \pm 1.74$	27.5 ± 5.6**	$1.0 \pm 0.3$
	10.2	9.6 ± 3.5**	42.8 ± 9.8**	$1.0 \pm 0.2$
	30.6	12.1 ± 2.7*	69.6 ± 13.2*	$0.9 \pm 0.1$
$PROD^b$	0	$0.4 \pm 0.2$	$21 \pm 7.2$	$11.2 \pm 4.2$
	1.7	3.5 ± 1.2**	108 ± 12.4**	$12.3 \pm 3.1$
	10.2	18.9 ± 5.5*	219 ± 48.2*	$13.2 \pm 2.9$
	30.6	$22.2 \pm 6.7*$	202 ± 42.8*	$11.7 \pm 4.6$
Cyp3a1 <sup>a</sup>	0	$1.5 \pm 1.1$	$1.1 \pm 0.8$	$0.5 \pm 0.1$
<b>71</b>	1.7	$1.0 \pm 0.4$	$1.2 \pm 0.8$	$0.6 \pm 0.1$
	10.2	$2.4 \pm 1.0$	$7.6 \pm 2.6**$	$0.7 \pm 0.1$
	30.6	3.7 ± 2.2**	16.3 ± 5.8*	$0.6 \pm 0.1$
$BROD^b$	0	15.1 ± 11.2	$52.1 \pm 22.8$	$25.3 \pm 10.1$
	1.7	$35.0 \pm 10.7$	$301.0 \pm 100.7**$	$25.9 \pm 11.2$
	10.2	148.3 ± 42.0**	1818.4 ± 420.1*	$26.2 \pm 12.8$
	30.6	201.1 ± 82.5*	2120.5 ± 582.5*	$26.3 \pm 11.1$

*Note*. Phase I hepatic microsomal EROD, PROD, and BROD activity along with real-time PCR of Cyp1a1, Cyp2b1, Cyp2b2, and Cyp3a1 gene expression were measured. Microsomes and RNA were isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

<sup>a</sup>Data are expressed as relative quantitation mean ± SD.

<sup>b</sup>Data are expressed as pmol/mg protein/min  $\pm$  SD, n = 5.

expression increased at the 1.7, 10.2, and 30.6 mg/kg/day doses by 4-, 13-, and 20.2-fold at PND 4, and 6.8-, 16.3-, and 26.8-fold at PND 21, respectively. Hepatic Cyp2b2 mRNA expression also increased significantly at the 1.7, 10.2, and 30.6 mg/kg/day doses by 1.9-, 5.6-, and 7-fold at PND 4, and 2.7-, 6.6-, and 10.6-fold at PND 21, respectively. In agreement, PROD showed a significant interaction of age and dose ( $F_{6.48} = 33.53$ ; p < 0.0001) with at all doses tested at PND 4 and PND 21, but not PND 60. Hepatic PROD, a marker for overall CYP2B activity, increased at the 1.7, 10.2, and 30.6 mg/kg/day doses by 8.8-, 47.2-, and 55.5-fold at PND 4, and 5.1-, 10.4-, and 9.6-fold at PND 21, respectively.

Cyp3a1 mRNA expression and BROD activity were used as markers for PXR receptor activation (Xie *et al.*, 2000). There was a significant interaction of age and dose for Cyp3a1

TABLE 3
Effect of DE-71 on Hepatic UGT-T4 Protein Activity and
Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
UGT-T <sub>4</sub> <sup>a</sup>	0	$0.6 \pm 0.1$	$0.6 \pm 0.1$	$0.5 \pm 0.1$
•	1.7	$0.6 \pm 0.1$	$0.6 \pm 0.1$	$0.5 \pm 0.0$
	10.2	$0.7 \pm 0.1$	$0.7 \pm 0.2$	$0.5 \pm 0.1$
	30.6	1.5 ± 0.3**	1.7 ± 0.4**	$0.5 \pm 0.1$
Ugt1a1 <sup>b</sup>	0	$1.2 \pm 0.4$	$0.2 \pm 0.1$	$0.1 \pm 0.1$
C	1.7	$1.7 \pm 0.5$	$0.2 \pm 0.1$	$0.1 \pm 0.0$
	10.2	$1.7 \pm 0.4$	$0.2 \pm 0.0$	$0.1 \pm 0.0$
	30.6	$1.5 \pm 0.2$	$0.1 \pm 0.0$	$0.1 \pm 0.0$
Ugt1a6 <sup>b</sup>	0	$1.8 \pm 1.0$	229.4 ± 12.1	$79.7 \pm 16.0$
C	1.7	$3.4 \pm 1.3$	$245.2 \pm 34.2$	$98.3 \pm 16.2$
	10.2	12.8 ± 4.8**	$229.5 \pm 17.8$	$108.7 \pm 16.9$
	30.6	34.0 ± 10.5*	$201.1 \pm 30.2$	$112.1 \pm 17.8$
Ugt1a7 <sup>b</sup>	0	$1.8 \pm 0.2$	$0.3 \pm 0.1$	$0.003 \pm 0.0003$
C	1.7	$2.5 \pm 0.5$	$0.3 \pm 0.1$	$0.003 \pm 0.0005$
	10.2	2.9 ± 0.8**	$0.4 \pm 0.1$	$0.003 \pm 0.0002$
	30.6	3.8 ± 1.5**	$0.7 \pm 0.2*$	$0.004 \pm 0.0005$
Ugt2b <sup>b</sup>	0	$1.2 \pm 0.4$	$1.2 \pm 0.7$	$7534 \pm 2963$
0	1.7	$1.0 \pm 0.4$	$7.6 \pm 6.0$	$10,043 \pm 3581$
	10.2	$1.3 \pm 1.0$	$1.5 \pm 0.3$	$9881 \pm 2274$
	30.6	$6.5 \pm 2.2*$	$78.0 \pm 9.3*$	$11,318 \pm 2227$

*Note.* Phase II hepatic microsomal enzyme activity was measured for UGTs using  $T_4$  as a substrate along with real-time PCR of Ugt1a1, Ugt1a6, Ugt1a7, and Ugt2b gene expression. Microsomes and RNA were isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

( $F_{6,48} = 17.66$ , p < 0.0001). A post hoc test indicated a significant effect at all doses tested at PND 4 and PND 21, but not at PND 60. Hepatic Cyp3a1 mRNA expression increased with 30.6 mg/kg/day treatment by 2.6-fold at PND 4 and 6.9 and 14.7-fold at PND 21 with 10.2 and 30.6 mg/kg/day treatment, respectively. In agreement with this, a significant interaction of age and dose ( $F_{6,48} = 64.90$ ; p < 0.0001) was observed for BROD at all doses tested at PND 4 and 21. BROD increased 9.8- and 13.3-fold at PND 4 with treatments of 10.2 and 30.6 mg/kg/day, and 5.8-, 34.9-, and 40.7-fold at PND 21 with treatments of 1.7, 10.2, and 30.6 mg/kg/day, respectively.

## UGTs: Enzyme Activity and Gene Expression

A significant interaction of age and dose ( $F_{6,48} = 12.81$ ; p < 0.0001) was detected for UGT-T<sub>4</sub> enzyme activity (Table 3). A post hoc test indicated a significant effect of DE-71 at the highest dose with a 2.5- and 2.8-fold increase at PND 4 and 21, respectively. No induction at PND 60 was observed. To determine which UGT isoform(s) may contribute to the overall enzyme activity, hepatic UGT mRNA expressions were further examined.

For Ugt1a1 mRNA expression, a significant interaction of age  $(F_{6,48} = 4.755; p < 0.0001)$  but not dose was observed at every dose group. However, a significant interaction of age and dose for Ugt1a6 mRNA expression ( $F_{6,48} = 4.755$ ; p < 0.0001) was identified at 10.6 and 30.6 mg/kg/day, with an increase of 7.2- and 19.6-fold at PND 4, respectively. Additionally, there was a significant interaction of age and dose for Ugt1a7 mRNA expression ( $F_{6,48} = 3.479$ ; p < 0.0062) with an increase of 1.6and 2.1-fold at 10.6 and 30 mg/kg/day, respectively, on PND 4 and 2.6-fold at the high dose on PND 21. Lastly, a significant effect of age for Ugt2b mRNA expression ( $F_{6.48} = 10.34$ ; p <0.0001) was identified. The large increase in developmental expression observed for Ugt2b may possibly mask a dose effect; therefore a one-way ANOVA was performed which identified a significant increase with 30.6 mg/kg/day treatment of 5.5- and 67.6-fold at PND 4 and 21, respectively.

Thus, although a significant induction of gene expression was observed at the middle dose, there is no concomitant increase in UGT-T<sub>4</sub> activity suggesting the enzyme assay used may not be sensitive enough to reflect the observed changes in gene expression.

# SULTs: Enzyme Activity and Gene Expression

Exposure to DE-71 led to a significant effect on age for SULT-T<sub>4</sub> ( $F_{6,48} = 11.25$ ; p < 0.0001), Sult1a1 ( $F_{6,48} = 154.4$ ; p < 0.0001) and Sult1c1 ( $F_{6,48} = 31.39$ ; p < 0.0001) with no dose related changes (Table 4). However, a significant interaction of age and dose was identified for Sult1b1 mRNA expression ( $F_{6,48} = 7.540$ ; p < 0.0001). A *post hoc* test identified a 2.8-fold increase on PND 4 with exposure to the high dose, and 2.1- and 4.1-fold increase at 10.2 and 30.6 mg/kg/day on PND 21, respectively.

#### Efflux Transporters: Gene Expression

A significant interaction between age and dose for the efflux transporter Mdr1 ( $F_{6,48} = 10.34$ ; p < 0.0001) was observed (Table 5). A post hoc test indicated an effect of DE-71 at all doses for PND 4 and 21, but not at PND 60. There was a dosedependent increase in hepatic Mdr1 mRNA expression of 1.5-, 1.8-, and 2.4-fold at PND 4 and 1.5-, 1.8-, and 2.2-fold at PND 21. Similarly, a significant interaction of age and dose for Mrp2 mRNA expression ( $F_{6,48} = 27.65$ ; p < 0.0001) was also observed. Mrp2 mRNA gene expression levels increased 1.6fold at the 30.6 mg/kg/day treatment group for PND 4. The expression of Mrp2 at PND 21 increased 1.5-, 1.9-, and 3.4fold, respectively. At PND 60, a significant 30 and 40% reduction at 10.2 and 30.6 mg/kg/day was also found. Lastly, a significant interaction of age and dose was observed for Mrp3 mRNA expression ( $F_{6,48} = 17.62$ ; p < 0.0001). A post hoc test indicated an effect for all doses at PND 4 and 21, but not at PND 60. The expression of hepatic Mrp3 mRNA, a major sinusoidal efflux transporter of glucuronides, indicated a 1.6-, 2.5-, and 4.1-fold increase at PND 4 and a 2.2-, 2.8-, and 5.1fold increase at PND 21.

<sup>&</sup>lt;sup>a</sup>Data are expressed as pmol/mg protein/min  $\pm$  SD, n = 5.

<sup>&</sup>lt;sup>b</sup>Data are expressed as relative quantitation mean  $\pm$  SD.

TABLE 4
Effect of DE-71 on Hepatic SULT-T4 Protein Activity and
Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60		
SULT-T <sub>4</sub> <sup>a</sup>	0	$0.2 \pm 0.2$	$0.6 \pm 0.1$	$0.5 \pm 0.2$		
	1.7	$0.1 \pm 0.0$	$0.4 \pm 0.3$	$0.5 \pm 0.4$		
	10.2	$0.1 \pm 0.1$	$0.4 \pm 0.2$	$0.6 \pm 0.4$		
	30.6	$0.3 \pm 0.2$	$0.5 \pm 0.3$	$0.6 \pm 0.4$		
Sult1a1 <sup>b</sup>	0	$1.0 \pm 0.3$	$2.2 \pm 0.2$	$0.34 \pm 0.1$		
	1.7	$1.2 \pm 0.3$	$2.7 \pm 0.5$	$0.35 \pm 0.1$		
	10.2	$1.1 \pm 0.3$	$2.1 \pm 0.5$	$0.40 \pm 0.1$		
	30.6	$1.1 \pm 0.6$	$2.2 \pm 0.6$	$0.39 \pm 0.2$		
Sult1b1 <sup>b</sup>	0	$1.1 \pm 0.1$	$3.0 \pm 0.6$	$2.4 \pm 0.6$		
	1.7	$1.8 \pm 0.1$	$3.6 \pm 0.2$	$2.1 \pm 0.8$		
	10.2	$1.1 \pm 0.2$	$6.4 \pm 0.8**$	$2.0 \pm 0.5$		
	30.6	$2.9 \pm 0.6**$	12.4 ± 1.0**	$2.0 \pm 0.1$		
Sult1c1 <sup>b</sup>	0	$1.7 \pm 0.6$	$4.1 \pm 1.5$	$1.0 \pm 0.1$		
	1.7	$2.1 \pm 0.5$	$5.0 \pm 2.5$	$1.0 \pm 0.2$		
	10.2	$2.3 \pm 0.9$	$3.8 \pm 0.8$	$1.1 \pm 0.5$		
	30.6	$2.2 \pm 0.4$	$4.5 \pm 3.3$	$0.9 \pm 0.4$		

*Note*. Phase II hepatic sulfotransferase enzyme activity was measured using  $T_4$  as a substrate along with real-time PCR of Sult1a1, Sult1b1, and Sult1c1 gene expression. Cytosolic fractions and RNA were isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

Influx Transporters: Gene Expression

A significant effect of age was observed for Oat2 ( $F_{6,48} = 270.5$ ; p < 0.0001) and Ntcp ( $F_{6,48} = 42.79$ ; p < 0.0001) but no dose effect was present (Table 6). However, a significant

TABLE 5
Effect of DE-71 on Hepatic Efflux Transporter Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
Mdr1 <sup>a</sup>	0	1.1 ± 0.1	$2.3 \pm 0.1$	2.4. ± 0.2
	1.7	$1.7 \pm 0.1**$	$3.3 \pm 0.4**$	$2.3 \pm 0.3$
	10.2	$1.9 \pm 0.8**$	$4.1 \pm 0.4**$	$2.7 \pm 0.0$
	30.6	$2.6 \pm 0.1**$	$5.0 \pm 0.9**$	$2.3 \pm 0.9$
$Mrp2^a$	0	$0.8 \pm 0.2$	$1.1 \pm 0.1$	$1.0 \pm 0.1$
-	1.7	$0.9 \pm 0.2$	$1.6 \pm 0.2**$	$1.0 \pm 0.1$
	10.2	$0.9 \pm 0.1$	$2.0 \pm 0.2**$	$0.7 \pm 0.0**$
	30.6	$1.3 \pm 0.0**$	$2.8 \pm 0.9**$	$0.6 \pm 0.0**$
$Mrp3^a$	0	$1.3 \pm 0.2$	$2.4 \pm 0.3$	$4.24 \pm 0.4$
_	1.7	$2.1 \pm 0.0**$	$5.2 \pm 1.1**$	$4.11 \pm 0.6$
	10.2	$3.3 \pm 1.7**$	$6.6 \pm 1.6**$	$3.83 \pm 0.3$
	30.6	5.1 ± 1.1**	$12.2 \pm 2.3*$	$4.14 \pm 0.9$

*Note.* Phase III hepatic efflux transporters were measured for Mdr1, Mrp2, and Mrp3 by real-time PCR. RNA was isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

TABLE 6
Effect of DE-71 on Hepatic Influx Transporter Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
Oat2 <sup>a</sup>	0	$1.3 \pm 0.3$	9.5 ± 1.3	9.5 ± 0.5
	1.7	$1.2 \pm 0.1$	$8.4 \pm 1.7$	8.5 ± 1.9
	10.2	$1.5 \pm 0.3$	$8.5 \pm 1.0$	$9.2 \pm 1.6$
	30.6	$1.1 \pm 0.2$	$7.7 \pm 2.0$	$9.0 \pm 0.7$
Oatp1a4 <sup>a</sup>	0	$1.1 \pm 0.1$	$53.1 \pm 2.1$	$40.0 \pm 5.5$
	1.7	$1.2 \pm 0.0$	$54.0 \pm 2.2$	$43.5 \pm 4.4$
	10.2	$1.2 \pm 0.1$	$57.4 \pm 2.5$	$40.0 \pm 3.8$
	30.6	$1.3 \pm 0.1$	69.4 ± 5.2**	$42.9 \pm 6.3$
Ntcp <sup>a</sup>	0	$1.2 \pm 0.1$	$0.8 \pm 0.1$	$1.0 \pm 0.2$
	1.7	$1.2 \pm 0.1$	$0.8 \pm 0.1$	$1.0 \pm 0.2$
	10.2	$1.2 \pm 0.1$	$0.8 \pm 0.1$	$1.0 \pm 0.1$
	30.6	$1.1 \pm 0.2$	$0.8 \pm 0.1$	$1.0 \pm 0.0$

*Note*. Phase III hepatic influx transporters were measured for Oat2, Oatp1a4, and Ntcp by real-time PCR. RNA was isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

interaction of age and dose for Oatp1a4 mRNA expression exists ( $F_{6,48} = 7.540$ ; p < 0.0001). A *post hoc* test identified a 1.3-fold increase for 30.6 mg/kg/day at PND 21. No effect at PND 4 and 60 were observed for Oatp1a4.

Transthyretin: Gene Expression

There was a significant effect of age ( $F_{6,48} = 205.7$ ; p < 0.0001) and dose ( $F_{6,48} = 3.02$ ; p < 0.039) observed for Ttr (Table 7). A *post hoc* test indicated that Ttr mRNA expression was significantly decreased 20% at the highest dose tested, 30.6 mg/kg/day, on PND 21. No effects were seen at PND 4 or 60.

Deiodinase 1: Enzyme Activity and Gene Expression

D1 is a marker for acute changes in TH metabolism (Zoeller *et al.*, 2006) (Table 8). A significant interaction of age and dose

TABLE 7
Effect of DE-71 on Hepatic Transthyretin Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
Ttr <sup>a</sup>	0	$1.0 \pm 0.1$	$1.0 \pm 0.1$	$0.4 \pm 0.0$
	1.7	$1.0 \pm 0.0$	$1.0 \pm 0.1$	$0.4 \pm 0.0$
	10.2	$0.9 \pm 0.2$	$0.9 \pm 0.1$	$0.4 \pm 0.1$
	30.6	$0.9 \pm 0.2$	$0.8 \pm 0.2**$	$0.5 \pm 0.0$

*Note*. The serum binding protein, transthyretin, was measured by real-time PCR. RNA was isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21. Significantly different from control: (p < 0.01) \* or (p < 0.05)\*\*.

<sup>&</sup>lt;sup>a</sup>Data are expressed as pmol/mg protein/min  $\pm$  SD, n = 5.

<sup>&</sup>lt;sup>b</sup>Data are expressed as relative quantitation mean ± SD.

<sup>&</sup>lt;sup>a</sup>Data are expressed as relative quantitation mean  $\pm$  SD, n = 5.

<sup>&</sup>lt;sup>a</sup>Data are expressed as relative quantitation mean  $\pm$  SD, n = 5.

<sup>&</sup>lt;sup>a</sup>Data are expressed as relative quantitation mean  $\pm$  SD, n = 5.

TABLE 8
Effect of DE-71 on Hepatic Deiodinase 1 Protein Activity and
Gene Expression

Target	Treatment (mg/kg/day)	PND 4	PND 21	PND 60
D1 <sup>a</sup>	0	257 ± 21	402 ± 99	500 ± 47
	1.7	111 ± 11*	$312 \pm 30$	$472 \pm 56$
	10.2	102 ± 10*	$347 \pm 32$	$489 \pm 37$
$d1^b$	30.6	$114 \pm 23*$	$115 \pm 19*$	$492 \pm 29$
	0	$1.7 \pm 0.4$	$9.0 \pm 0.9$	$0.5 \pm 0.1$
	1.7	$1.6 \pm 0.2$	$8.7 \pm 0.6$	$0.4 \pm 0.1$
	10.2	$1.1 \pm 0.1**$	$8.3 \pm 1.6$	$0.5 \pm 0.2$
	30.6	$0.9 \pm 0.3*$	5.9 ± 1.3*	$0.5 \pm 0.1$

*Note.* Deiodinase 1 enzyme activity was measured using  $T_4$  as a substrate along with gene expression by real-time PCR. RNA was isolated from male rat livers at PND 4, 21, and 60 after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 (0, 1.7, 10.2, and 30.6 mg/kg/day) from GD 6 to PND 21.

Significantly different from control:  $(p < 0.01)^*$  or  $(p < 0.05)^{**}$ .

for D1 enzyme activity ( $F_{6,48}=15.02$ ; p<0.0001) was observed for all doses tested at PND 4 and for 30.6 mg/kg/day at PND 21. At PND 4, D1 activity decreased 60% at all doses, whereas at PND 21, a significant 70% decrease in activity was observed only at the highest dose. There was also a significant interaction of age and dose for d1 mRNA expression ( $F_{6,48}=5.978$ ; p<0.0001). A *post hoc* test indicated an effect at doses of 10.2 and 30.6 mg/kg/day for PND 4 and at 30.6 mg/kg/day for PND 21, but not at PND 60. At PND 4, expression levels were decreased in a dose related fashion to 40 and 50% at 10.2 and 30.6 mg/kg/day, respectively. However, on PND 21, the only decrease in d1 expression was at the high dose. No changes were observed at PND 60.

### DISCUSSION

Previous hypotheses on perinatal TH disruption by DE-71 have focused on induction of hepatic UGT-T<sub>4</sub>-mediated TH catabolism resulting in decreased circulating T<sub>4</sub> levels. It has also been hypothesized that competitive binding between PBDE metabolites and TH to serum transport proteins affects thyroid homeostasis. This study further investigates parameters involved in the alteration of TH levels with a focus on nuclear receptor-mediated activation during development, in the presence of a PBDE mixture. Specifically, this study aims to further identify genes activated by AhR, CAR, and PXR during perinatal exposure to DE-71. This information will further the risk assessment and provide information as to the mechanisms by which this commercial penta mixture exerts its effects. Concomitant with a decrease in T<sub>4</sub>, we observed an increase in hepatic UGT-T<sub>4</sub> activity and UGT mRNA isozyme gene

expression, a decrease in D1 enzyme activity and mRNA expression, and increases in the transcription of Sult1b1 and transporters of TH or glucuronides in neonatal and juvenile male rats following maternal PBDE treatment.

After perinatal exposure to DE-71, a dose-dependent induction of CYP1A1 and CYP2B demonstrates a similar induction pattern as previously reported (Zhou et al., 2002), whereas we observe for the first time a dose-dependent induction of CYP3A. This is in agreement with a 28-day oral exposure study using a purified DE-71 mixture in adult Wistar rats (van der Ven et al., 2008). This is also the first report measuring the induction of CYP450 mRNA expression after perinatal exposure to DE-71. The induction of CYP1A1, CYP2B, and CYP3A along with their respective mRNA transcripts signifies DE-71 is an agonist for AhR, CAR, and PXR, respectively. The increased induction seen in CYP2B as compared with CYP3A supports the suggestion that PBDE congeners have a preference for CAR over PXR (Pacyniak et al., 2007; Sanders et al., 2005; Richardson et al., 2008). Although individual PBDE congeners have been shown not be AhR agonists (Peters et al., 2006), induction of CYP1A1 gene expression and protein further supports the presence of dioxinlike contaminants in the DE-71 mixture (Hanari et al., 2006) as being responsible for the Ah receptor-based responses (Sanders et al., 2005).

Our data correlate with previous UGT developmental enzyme activity studies in Long-Evans male rats which also demonstrated decreases in TH concentrations along with increases in hepatic UGT-T<sub>4</sub> activity with exposure to DE-71 (Zhou et al., 2002). Different UGT isoforms have unique developmental patterns and their regulation is thought to be nuclear receptor specific. In this study, perinatal DE-71 exposure increased AhR mediated Ugt1a6 (Auyeung et al., 2003; Nishimura et al., 2005) and Ugt1a7 (Metz and Ritter, 1998) along with CAR mediated Ugt2b (Zhou et al. 2005) mRNA expression. Members of the UGT1A family specifically glucuronidate T<sub>4</sub>, whereas members of the UGT2b family glucuronidate T<sub>3</sub> (Vansell and Klaassen, 2002). The significant increase in hepatic Ugt1a6 and Ugt1a7 mRNA expression further supports contaminants in DE-71 as agonists for the AhR. In addition, PBDE induction of Ugt2b at the high dose parallels the UGT induction pattern and is likely CAR mediated.

In this study, there is a lack of DE-71 effect on SULT-T<sub>4</sub>, Sult1a1, and Sult1c1, whereas an increase in Sult1b1 mRNA expression was identified. SULT1A1, 1B1, and 1C1 have enzymatic activity towards T<sub>3</sub> and T<sub>4</sub>, whereas SULT1C1 exhibits higher expression in males. The lack of a sex difference in Sult1b1 expression suggests it has a similar role in both sexes. For this reason it has been hypothesized that SULT1B1 may be more important for TH homeostasis than other isoforms (Dunn *et al.*, 1999; Fujita *et al.*, 1997). Considering sulfation is a reversible pathway of TH metabolism which depends on the free hormone recovery by sulfatases (Darras *et al.*, 1999), specific induction of Sult1b1 may be

<sup>&</sup>lt;sup>a</sup>Data are expressed as pmol/mg protein/min  $\pm$  SD, n = 5.

<sup>&</sup>lt;sup>b</sup>Data are expressed as relative quantitation mean ± SD.

important in TH homeostasis during exposure to endocrine disrupting compounds such as PBDEs. *In vitro* rodent experiments have demonstrated hepatic SULTs having different conjugation affinities for iodothyronines with  $T_4 < rT_3 < T_3 < T_2$  (Kaptein *et al.*, 1997). Although the Km of rodent SULT- $T_4$  is high in relation to other iodothyronines,  $T_4$  comprises the fractional majority of TH at any given time in circulation and can possibly contribute to a large fraction of  $T_4$  for sulfation. The lack of effect observed on hepatic SULT- $T_4$  activity after exposure to DE-71 contrasts with the observed increases in Sult1b1 mRNA expression. This could be due to SULT- $T_4$  enzyme activity assay not evaluating the activity of specific SULT isoforms. Alternatively, increases in Sult mRNA expression may not result in marked increases of the respective enzymes.

The efflux transporters Mdr1, Mrp2, and Mrp3 are members of the ABC binding cassette superfamily (Borst and Elferink, 2002; Dean and Allikmets, 2001) and are regulated by AhR, CAR and PXR (Cherrington et al., 2002; Geick et al., 2001; Johnson et al., 2002; Kast et al., 2002; Maglich et al., 2002; Teng et al., 2003; Xiong et al., 2002). Mrp2 resides at the canalicular membrane and secretes its substrates into bile (Müller et al., 1996). Mrp3 is present at low levels at the basolateral membrane for export of substrates into sinusoidal blood. The expression of Mdr1 and Mrp2 are inducible by hormones and steroids and their activity may change during development (Courtois et al., 1999; Demeule et al., 1999). The induction of Mdr1 and Mrp2 both occurred at PND 4 and 21 with Mdr1 having a greater sensitivity at PND 4. Although Mrp2 and Mdr1 both have a high sensitivity to PBDEs later in development, Mdr1 efflux mechanisms appear to be involved in elimination and detoxification during early postnatal development to a greater extent than Mrp2. The developmental expression of the basolateral efflux transporter Mrp3 has been previously documented in mice (Maher et al., 2005) and correlates with our findings in rats. Its age and dose-dependent sensitivity to PBDEs appear to be similar to the canalicular efflux transporter Mdr1; however Mrp3 is induced to a greater

In rat liver, basolateral uptake systems include the sodium taurocholate cotransport protein (NTCP) and the OATPs. The basolateral Na<sup>+</sup>-dependent bile salt transporter, NTCP, is specific to hepatocytes and is distributed homogeneously throughout the liver (Stieger et al., 1994). Bile salts are the major substrate for NTCP, however, other compounds such as estrogen conjugates, TH, and xenobiotics that are covalently bound to taurocholate can also be transported (Kouzuki et al., 2000). DE-71 did not alter mRNA expression levels of Ntcp. The lack of effect seen with exposure to this commercial mixture implies Ntcp is not activated by AhR/CAR/PXR. In addition, full maturation of NTCP transport activity is delayed until 4 weeks of age due to incomplete glycosylation (Kühlkamp et al., 2005). This delay may contribute to the lack of Ntcp mRNA expression alterations during perinatal exposure seen here.

TABLE 9
Summary on the Effect of DE-71 on THs, Hepatic Protein Activity, and Gene Expression<sup>a</sup>

	3,		
Target	PND 4	PND 21	PND 60
$T_4$	$\downarrow \downarrow$	$\downarrow\downarrow$	_
$T_3$	_	_	_
EROD	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	_
PROD	$\uparrow\uparrow\uparrow$	$\uparrow \uparrow$	_
BROD	$\uparrow \uparrow$	$\uparrow\uparrow\uparrow$	_
UGT-T <sub>4</sub>	$\uparrow$	<b>↑</b>	_
SULT-T <sub>4</sub>	_	_	_
D1-T4	$\downarrow\downarrow\downarrow$	$\downarrow$	_
Cyp1a1	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	$\uparrow$
Cyp2b1	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	_
Cyp2b2	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	_
Cyp3a1	<b>↑</b>	$\uparrow \uparrow$	_
Ugt1a1	_	_	_
Ugt1a6	$\uparrow \uparrow$	_	_
Ugt1a7	<b>↑</b>	<b>↑</b>	_
Ugt2b	$\uparrow$	$\uparrow \uparrow$	_
Sult1a1	_	_	_
Sult1b1	_	$\uparrow \uparrow$	_
Sult1c1	_	_	_
Mdr1	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	
Mrp2	$\uparrow$	$\uparrow\uparrow\uparrow$	<u> </u>
Mrp3	$\uparrow\uparrow\uparrow$	$\uparrow\uparrow\uparrow$	_
Oatp1a4	_	<b>↑</b>	_
Oat2	_	_	_
Ntcp	_	_	_
Ttr	_	$\downarrow$	_
d1	$\downarrow \downarrow$	$\downarrow$	_

*Note.* Overall hepatic effects after perinatal exposure to the polybrominated diphenyl ether mixture, DE-71 on THs, hepatic protein activity and gene expression. Data was measured in male rat at PND 4, 21, and 60 after perinatal exposure to 0, 1.7, 10.2, and 30.6 mg/kg/day between GD 6 to PND 21.

<sup>a</sup>Three, two, and one arrow refers to significant effects at doses 1.7, 10.2, and 30.6 mg/kg/day, respectively.

Although NTCP represents the major hepatocellular uptake system for conjugated bile salts, OATPs mediate sodium-independent uptake of a large variety of substrates. Specifically, OATP1a4 actively transports bile acids, xenobiotics, and TH. OATP1a4 belongs to the ABC cassette superfamily and is regulated by CAR and PXR (Wagner *et al.*, 2005). Because OATP1a4 is a basolateral hepatic influx transporter, the increased expression levels seen here may indicate a demand for sequestration of PBDEs and T<sub>4</sub> into the liver for biotransformation and elimination.

Oat2 is found in higher concentrations in adult male rat liver than in kidney, however, the opposite is true for females (Buist *et al.*, 2002; Pavlova *et al.*, 2000). In this study, the expression levels of Oat2 in male Long-Evans rats were low at PND 4 and increased at PND 21, and then remained high through PND 60. This developmental pattern is consistent with previous findings for rat Oat2 in the liver (Simonson *et al.*, 1994). The level of transcript increases

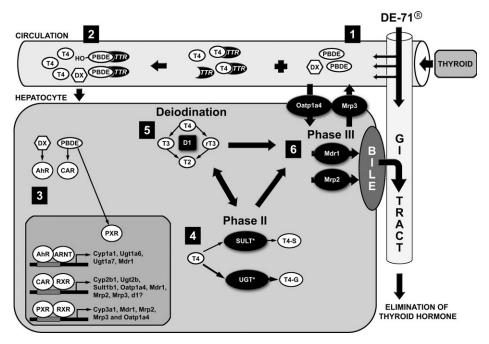


FIG. 1. Possible mechanisms of TH disruption after perinatal exposure to the commercial "penta" PBDE flame-retardant mixture, DE-71. (1) PBDE congeners and dioxin-like (DX) contaminants enter the circulation. (2) PBDE (parent or hydroxylated metabolite) displace  $T_4$  from its serum binding protein, transthyretin (TTR). This result in an increase of free  $T_4$  destined for hepatic metabolism and elimination. (3) PBDE and DX activate nuclear receptors initiating transcription of XMEs. (4) XMEs consequently conjugate  $T_4$  by phase II enzymes, UGT and SULT. (5) Deiodinase 1 (D1) can deiodinate parent and conjugated  $T_4$  along with their metabolites. (6) Influx transporters (Oatp1a4) further increase the uptake of  $T_4$  for metabolism. Efflux transporters eliminate  $T_4$  or its conjugates from hepatocytes either in the serum (Mrp3) or the bile (Mdr1 and Mrp2).

dramatically within two days post-partum and continues until the level stabilizes within the second week of postnatal development.

D1 is mainly expressed in the liver, thyroid, and kidney. Evidence suggests D1 is regulated directly or indirectly by CAR (Tien et al., 2007) however time and dose response relationship observed here suggests D1 is regulated by CAR and/or AhR but not PXR. Raasmaja et al. (1996) suggest a possible mechanism for the reduction in T<sub>4</sub> to be due to increased tissue-specific deiodinase activity converting T<sub>4</sub> to T<sub>3</sub>. In this study, hepatic D1 activity decreased 60% at all doses on PND 4, and 70% with the highest dose on PND 21. Similar decreased serum T<sub>4</sub> and hepatic D1 enzyme activity were seen with the PCB mixture, Aroclor 1254 (Hood and Klaassen, 2000). Considering the similarities in PCB and PBDE structures and effects, common mechanisms are likely involved in D1 reduction. Interestingly, decreases in rat hepatic D1 have also been observed following exposure to 2,3,7,8-tetrachlorodibenzop-dioxin (TCDD) and suggests D1 effects seen here may be attributed, at least in part, to the dioxin-like contaminants (Viluksela et al., 2004). However, decreases in D1 were a secondary effect to circulating T4 reduction after TCDD exposure. Future studies are needed to determine whether the sensitive decrease in D1 activity is also observed with exposure to purified PBDE congeners.

The reduction in D1 activity and mRNA message levels after perinatal exposure to the penta PBDE commercial mixture has proven challenging to explain. D1 is thought to be responsible for the major portion of  $T_3$  production peripherally. However, during hypothyroidism, plasma  $T_4$  is reduced and peripheral  $T_3$  conversion from  $T_4$  is believed to be sustained by upregulation of extrahepatic D2 and downregulation of D1 (Zavacki *et al.*, 2005). D2 is a more catalytically efficient enzyme, maintaining  $T_3$  levels by increasing the fractional conversion of  $T_4$  to active  $T_3$  as compared with the equal conversion of active  $T_3$  and inactive  $T_3$  from D1.

In addition to D2, SULTs have been reported to work with D1 in the regulation of TH homeostasis. Sulfation in the human fetus has been proposed to be a protective mechanism regulating T<sub>3</sub> levels (Santini et al., 1993). This regulation is mediated by deiodinases (T<sub>4</sub>S to T<sub>3</sub>S), and furthermore by sulfatases (T<sub>3</sub>S to T<sub>3</sub>), during periods of high T<sub>3</sub> demands (Chopra, 1994; Santini et al., 1992). In this study, rat hepatic Sult1b1 was increased. It has been shown that deiodination rates by D1 of T<sub>4</sub> to inactive rT<sub>3</sub> are increased nearly 200 times by sulfation  $(T_4S \text{ to } rT_3S)$ , whereas deiodination of  $T_4$  to active T<sub>3</sub> is completely lost after sulfation (T<sub>4</sub>S to T<sub>3</sub>S) (Visser et al., 1993). Therefore, the decreases in rat hepatic D1 activity observed after exposure to xenobiotics may act as a protective mechanism to reduce the conversion of T<sub>4</sub>S to inactive rT<sub>3</sub>S. Favorably, this reduction of D1 serves to maintain T<sub>3</sub> levels.

Risk assessment of the PBDE commercial mixtures is ongoing and this *in vivo* study aids in a better evaluation of the

possible risks for human beings. Although the administered doses used here may seem high, this study as well as others, have shown that effects of PBDEs are seen in animal models at concentrations within 10-fold of the high end of the human population in North America (McDonald, 2005). This study demonstrates perinatal exposure as low as 1.7 mg DE-71/kg/ day is sufficient to alter hepatic enzyme activity measured as early as PND 4 (Table 9). The most sensitive endpoints in terms of both mRNA and enzyme activity were a reduction of D1 and induction of CYP1A1, 2B1/2, and 3A. These endpoints were all more sensitive at PND 4 than PND 21 with the exception of Cyp3a1/CYP3A. In contrast, Ugt2b, Sult1b1, and Mrp2 expression are all higher on PND 21 than PND 4, but these endpoints were less sensitive than CYP3A. Expression of Mdr1 and Mrp3 are also extremely sensitive at PND 4. In contrast, UGT mRNA effects were only seen at 10.2 mg/kg/ day or greater and enzyme activity at 30.6 mg/kg/day, indicating UGT-T<sub>4</sub> is not the most sensitive marker for this PBDE mixture. All effects were largely reversible by PND 60. The induction of hepatic Sult1b1 mRNA expression seen here along with decreases of D1 may work together to maintain serum T<sub>3</sub> and reduce T<sub>4</sub>. In addition, the hepatic efflux transporters Mdr1, Mrp2, and Mrp3 may be involved. However, studies to identify sulfated TH specific transporters and whether the alterations in mRNA levels seen in this study are reflected in protein levels and enzyme activity are clearly needed.

The  $T_4$  depleting effects of DE-71 are likely to involve multiple mechanisms of action including Phase II glucoronidation and sulfation, transthyretin displacement, decreased hepatic deiodinase 1 activity, and increases in hepatic transporter phase III elimination (Fig. 1). This study has demonstrated coordinate modification in the expression of transport proteins and detoxification enzymes in the postnatal period of development after low perinatal exposure to DE-71, a commercial PBDE mixture. These alterations could be responsible for the important and rapid changes in TH observed during this period of life. Because DE-71 contains both PBDEs and PBDDs/PBDFs, it is difficult to attribute every effect measured in this study to the sole activation of CAR/PXR or AhR, respectively. Considering household dust contains both PBDEs and dioxin-like compounds and is suspected to be a major source of exposure for humans and other animals, this commercial mixture study is important to the understanding of its real world toxic effects. In furthering the risk assessment of this commercial PBDE mixture, this data demonstrates that DE-71 disrupts TH homeostasis in rats during development via perinatal exposure; however, the mechanism(s) of action appear complex.

#### Supplementary Data

Supplementary data are available online at http://toxsci.oxfordjournals.org/.

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